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Range: from **begin** to **end** Features:  SNP  CDD  MGC  HPRD  :

**1: CAF33259. Reports unnamed protein p...[gi:45112868]** [BLink, Links](#)

LOCUS CAF33259 325 aa linear PLN 04-MAR-2004

DEFINITION unnamed protein product [Penicillium citrinum].

ACCESSION CAF33259

VERSION CAF33259.1 GI:45112868

DBSOURCE embl accession [CQ769141.1](#)

KEYWORDS .

SOURCE Penicillium citrinum

ORGANISM [Penicillium citrinum](#)  
 Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;  
 Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Penicillium.

REFERENCE 1

AUTHORS Asako, H. and Shimizu, M.

TITLE Modified reductase and its gene and use thereof

JOURNAL Patent: EP 1386961-A 04-FEB-2004;  
 Sumitomo Chemical Company, Limited (JP)

FEATURES Location/Qualifiers

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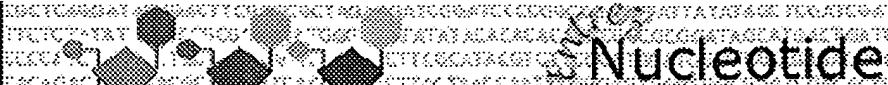
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Range: from  to   Reverse complemented strand Features:

1: [CQ769141](#). Report Sequence 2 from P...[gi:45112867] [Links](#)

**LOCUS** CQ769141 978 bp DNA linear PAT 04-MAR-2004  
**DEFINITION** Sequence 2 from Patent EP1386961.  
**ACCESSION** CQ769141  
**VERSION** CQ769141.1 GI:45112867  
**KEYWORDS**  
**SOURCE** Penicillium citrinum  
**ORGANISM** Penicillium citrinum  
 Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;  
 Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Penicillium.  
**REFERENCE** 1  
**AUTHORS** Asako, H. and Shimizu, M.  
**TITLE** Modified reductase and its gene and use thereof.  
**JOURNAL** Patent: EP 1386961-A 2 04-FEB-2004;  
 Sumitomo Chemical Company, Limited (JP)  
**FEATURES** Location/Qualifiers  
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**ORIGIN**

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Database : UniProt\_05.80:\*
   
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 2: uniprot\_trembl:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query				Description
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6	1191	68.8	327	2	Q6RZX1_TRIAT	Q6rzx1 trichoderma
7	1137.5	65.7	323	2	Q51QM9_MAGGR	Q51qm9 magnaporthe
8	1038	60.0	256	2	Q4I4F0_GIBZE	Q4i4f0 gibberella
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14	593.5	34.3	309	2	Q5KH94_CRYNE	Q5kh94 cryptococcu
15	583	33.7	310	2	Q6FR42_CANGA	Q6fr42 candida gla
16	575.5	33.2	310	2	Q6FY54_CANGA	Q6fy54 candida gla
17	574.5	33.2	312	1	GCY YEAST	P14065 saccharomyc
18	570.5	33.0	309	2	Q6CRC8_KLULA	Q6crc8 kluyveromyc
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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5	1731	100.0	325	8	ADK70247	Adk70247 Penicilli
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7	1731	100.0	325	8	ADM46567	Adm46567 Penicilli
8	1731	100.0	325	8	ADK51190	Adk51190 Wild-type
9	1731	100.0	325	8	ADN97092	Adn97092 3 hydroxy
10	1728	99.8	325	8	ADK51203	Adk51203 Mutant Pe
11	1726	99.7	325	8	ADK51204	Adk51204 Mutant Pe
12	1723	99.5	325	8	ADK51205	Adk51205 Mutant Pe
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15	574.5	33.2	312	2	AAW29217	Aaw29217 S. cerevi
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17	574.5	33.2	312	5	ABG93198	Abg93198 S. cerevi
18	574.5	33.2	312	8	ADS43942	Ads43942 Bacterial
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20	573.5	33.1	312	2	AAW29218	Aaw29218 S. cerevi
21	568	32.8	321	8	ADN19632	Adn19632 Bacterial

Not all  
same parents  
same date

RESULT 1  
 US-10-004-115B-1  
 ; Sequence 1, Application US/10004115B  
 ; Patent No. 6884607  
 ; GENERAL INFORMATION:  
 ; APPLICANT: ASAKO, HIROYUKI  
 ; APPLICANT: MATSUMURA, KENJI  
 ; APPLICANT: SHIMIZU, MASATOSHI  
 ; APPLICANT: ITO, NOBUYA  
 ; APPLICANT: WAKITA, RYUHEI  
 ; TITLE OF INVENTION: PROCESS FOR PRODUCING OPTICALLY ACTIVE  
 ; TITLE OF INVENTION: 4-HALO-3-HYDROXYBUTANOATE  
 ; FILE REFERENCE: 7372-72249  
 ; CURRENT APPLICATION NUMBER: US/10/004,115B  
 ; CURRENT FILING DATE: 2001-12-06  
 ; PRIOR APPLICATION NUMBER: JP 2000-372704  
 ; PRIOR FILING DATE: 2000-12-07  
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 ; PRIOR FILING DATE: 2001-01-15  
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 ; PRIOR FILING DATE: 2001-06-11  
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US-10-004-115B-1

parent  
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 ems19  
 n> variant

Query Match 100.0%; Score 1731; DB 2; Length 325;  
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Db

301 FRFVNMKDTFGYDVWPEETAKNLSA 325

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2: pir2:\*

3: pir3:\*

4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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9	529	30.6	290	2	T02543
10	527.5	30.5	309	2	B84599
11	523.5	30.2	294	2	T02542
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13	521	30.1	320	2	T48188
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